

RAW SEQUENCE LISTING

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Application Serial Number: 10/S19,811
Source: PCT
Date Processed by STIC: 1-12-05

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PCT

RAW SEQUENCE LISTING

DATE: 01/12/2005

PATENT APPLICATION: US/10/519,811

TIME: 09:45:17

Input Set : A:\892370 - seq.txt

Output Set: N:\CRF4\01122005\J519811.raw

5 <110> APPLICANT: University of Guelph
 7 <120> TITLE OF INVENTION: Novel Inducible Genes From Alfalfa And Method Of Use Thereof
 9 <130> FILE REFERENCE: 08-892370WO
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/519,811
 C--> 12 <141> CURRENT FILING DATE: 2004-12-28
 14 <150> PRIOR APPLICATION NUMBER: 60/392,444
 15 <151> PRIOR FILING DATE: 2002-06-28
 17 <160> NUMBER OF SEQ ID NOS: 19
 19 <170> SOFTWARE: PatentIn version 3.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 474
 23 <212> TYPE: DNA
 24 <213> ORGANISM: Medicago sativa
 26 <220> FEATURE:
 27 <223> OTHER INFORMATION: Nucleotide sequence of H7 coding region
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 34 gttgaaattg ttgaaggaaa tggaggaccc ggaactatta agaagctatc cattgttgaa 180
 36 gatggcaaaa ccaactttgt gctacacaaa ttagattcag tggatgaggc aaactttgga 240
 38 tataactaca gcttagtgagg aggaacaggg ttggatgaaa gtttagagaa agttgaattt 300
 40 gagacaaaaa ttgttgctgg ctctgatggg ggatccattg ttaagatttc agtgaaatac 360
 42 cataccaaag gtgatgcaac tctatctgaa gcagtacgtg aggagactaa ggccaaagga 420
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 52 <220> FEATURE:
 53 <223> OTHER INFORMATION: Nucleotide sequence of H11 coding region
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 60 ctaaagtttt cattagatca tcaaattaat atcaaacaaa cttctcttct atccctcaca 180
 62 gcaatcacat ttccattctt attggatacc aaagagtttg ggatatttga aggaagaaca 240
 64 tttgctctca ttcaccccat tgtgttgggt ggtttgttct tctatactct atatgctggc 300
 66 tatttggggg ggcaatggcg ccgagttagg actattcaaa atgatattaa tgagctcaag 360
 68 aaacaactca aacctgcacc ggctcgccct gatggtaaag cacttgaaac ttcaccgcca 420
 70 tcacctgttg aacttcaa at ccagaaactt actgaggaga ggaaagagct tatcaaaggt 480
 72 tcatcacagg ataaacactt taatgctgga tccatacttc taggatttgg tgtctttgag 540
 74 gctgttggtg tgaggactca acacatggtt aaggacagga aagctatttc caggtccaca 600
 76 tttatttgca ggagcaggca ttaccgtctt atgggcactg gcagcagctc tagtaccacc 660
 78 gatgcagaaa ggcagtga 678

(ps, 6)

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83 <212> TYPE: DNA
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86 <220> FEATURE:
87 <223> OTHER INFORMATION: Nucleotide sequence of H12 coding region
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92 aagagtcttt tacaaagtga tgctctttac cagtatatc tagagaccag tgtcttccca      120
94 agagaacatg aagccatgaa agagttgaga gaggtcacag caaacacccc atggaacatc      180
96 atgacaacct ctgcagatga aggacaattt ttgagcatgc tccttaaact tatcaatgct      240
98 aagaatacca tggaaattgg tgtctacact ggctactccc tccttgccac tgccttagct      300
100 attcctgaag atggaaagat tttggctatg gacattaaca aagaaaatta cgaattgggt      360
102 ctacctgtaa ttaaaaaagc tgggtttgat cacaaaattg atttcagaga aggtccagct      420
104 cttccagttc ttgatgaaat gatcaaagac gaaaagaatc atggtagcta cgatttcatt      480
106 tttgtggatg ctgacaaaga caattacctc aactaccata agaggttaat tgatcttggt      540
108 aaagtgggag gtgtgatcgg gtacgacaac accttatgga atggatctgt ggttgacccc      600
110 cctgatgctc cattgaggaa gtatgttagg tactatagag attttgttt ggagcttaac      660
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119 <212> TYPE: DNA
120 <213> ORGANISM: Medicago sativa
122 <220> FEATURE:
123 <223> OTHER INFORMATION: Nucleotide sequence of H7 regulatory region
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128 ccacccatga caccattgct atatttcaat ttgggaaaat attgctataa agttactgta      120
130 gtaactttta gaagaagggt ttttttttaa ggattttaga ggaaggtag caacacacat      180
132 gcactttaaa tatacathtt ttcttataaa gtttttgat cgagttgaga aatcatatat      240
134 atactcataa atcatgtgga ttcatataa tttaatagaa cacataaatt ttaaccgaga      300
136 aataaagtgt tgcaaatata tgttaaaaga gtacgttggt aacattattt taatttctt      360
138 tattcaatcc acactttgag tcatggactg ctatactaatt tcattttggt tttcgcaacc      420
140 taattagaga ttgtccagat acaaagagga gtaacctaat aaataaatat taaaatatc      480
142 accaacggcc tcagtaagct acttgagcta aacaatgaga tttccaaata aggtagggtcc      540
144 ttcccaagtt ctataaatag catccctcac catgtcataa accgcatcac aagttatata      600
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150 <211> LENGTH: 438
151 <212> TYPE: DNA
152 <213> ORGANISM: Medicago sativa
154 <220> FEATURE:
155 <223> OTHER INFORMATION: Nucleotide sequence of H11 regulatory region
157 <220> FEATURE:
158 <221> NAME/KEY: misc_feature
159 <222> LOCATION: (1)..(438)
160 <223> OTHER INFORMATION: where "n" is a or g or c or t or other
163 <400> SEQUENCE: 5

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166 cgcgtggtcg acggccccggg ctggtatcag cgagtaacga ttcattcatat ctccactag 120
168 ggatgaatga tttattattg agtttatgaa tttgaactat tacttctaata ttctaaatga 180
170 agacatttaa gtaaaagatt aaaatatctt agtttcaaat attttggtatt ttagaattta 240
172 aatttaattct ttaaaaaaaaa attaaattta aagaagataa aaaggagaa aataaataga 300
174 tgaatataat ttgtaaacat gaagacctta tctccagtaa aaaaacatat ggaccttatt 360
176 tttttgaggt aggaaggatc tacgcgggga acctcttctt gactgtgaac cccgtatgca 420
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186 <220> FEATURE:
187 <223> OTHER INFORMATION: Nucleotide sequence of H12 regulatory region
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194 aattttattat taaaattcac acttagatgg cctaaaaatt aacacttatt ttaacaatt 180
196 caaataaaat atacgacgaa atgagtgtaa tttagttggt taagcatcgt caaagcttgg 240
198 agagaaaagat catagtttga tctttgaaaa ctatactatt gaaaagggtg aagatatcta 300
200 acctccaaca aaattttatt gatagtcgat tcaaatattc aaaatttggg aaatatattg 360
202 taaattgtta agttgggaaa aatatgttaa ttttcaaatt accatttgca catttttcta 420
204 atctcaaatc acatttaagg gatgttgact actttcgttt tgtacaaatc tttacaattt 480
206 taacatttat aaaatgtgtt ttggtagata aaaagtgtga gtattcttta taagagattg 540
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210 attgtaagaa ttcattataa gattatgtca ttccctcaaa agaaaattag atgatgtcat 660
212 tttcataact cattttctat aaatacagaa aatcctcaaa aatgaaaaac ctcggtcaaa 720
214 aaataaaaga aaaacatcaa tagtggactg gccacactc attgctttgc tttagtagta 780
216 gaaagtagac ctcaccaacc acgaaccgga cgccgaccgg ttcaaccaa catcacacca 840
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224 <211> LENGTH: 1424
225 <212> TYPE: DNA
226 <213> ORGANISM: Medicago sativa
228 <220> FEATURE:
229 <223> OTHER INFORMATION: Nucleotide sequence of genomic H7
231 <400> SEQUENCE: 7
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234 ccacccatga caccattgct atatttcaat ttgggaaaat attgctataa agttactgta 120
236 gtaactttta gaagaagggt ttttttttaa ggattttaga ggaaggtag caacacacat 180
238 gcacttttaa tatacatttt ttcttataaa gtttttgat cgagttgaga aatcatatat 240
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242 aataaagtgt tgcaaatata tggtaaaaga gtacgttggt aacattattt taatttcttt 360
244 tattcaatcc acactttgag tcatggactg ctatactaat tcattttgtt tttcgcaacc 420
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256 cagctaaact ctacaaggct cttgcāaaag atgctgatga aatcgtccca aagggtgattt 780
258 ctgctgcccc aagtgttgaa attgttgaaag gaaatggagg acccggaact attaagaagc 840
260 tatccattgt tgaagatggc aaaaccaact ttgtgctaca caaattagat tcagtggatg 900
262 aggcaaaactt tggatataac tacagcttag tgggaggaac agggttggat gaaagttag 960
264 agaaagttga atttgagaca aaaattgttg ctggctctga tgggtggatcc attgttaaga 1020
266 tttcagtga ataccatacc aaaggtgatg caactctatc tgaagcagta cgtgaggaga 1080
268 ctaaggccaa aggaactgga cttatcaagg ccattgaggg ctacgtttta gcaaacctta 1140
270 attactagcc aattaaacc tattgaggac ttttaatttg gttgtgttgt ttcattgcga 1200
272 taataattaa agtttatgat gcgggttgaa tgtgttgagt atacatcaag gtctttggct 1260
274 cgtacatgtg tggtggcttt gttggatggt gtgaggtttg agtgctattt tgggtgttta 1320
276 aaaacaaaaa cctatgttgt gttgggtgata aggttttgca ccattctgtat tatgcaataa 1380
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281 <210> SEQ ID NO: 8

282 <211> LENGTH: 1482

283 <212> TYPE: DNA

284 <213> ORGANISM: Medicago sativa

286 <220> FEATURE:

287 <223> OTHER INFORMATION: Nucleotide sequence of genomic H11

289 <220> FEATURE:

290 <221> NAME/KEY: misc_feature

291 <222> LOCATION: (1)..(1482)

292 <223> OTHER INFORMATION: Where n is a or g or c or t or other

295 <400> SEQUENCE: 8

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316 tctcttctat ccctcacagc aatcacattt ccattcttat tggataccaa ggcaagcaag 660
318 caagcaagca tcctattcta ttctattctt tcatccatat ctttactctt ttgttttcta 720
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322 ggatatttga aggaagaaca tttgctctca ttcaccccat tgtgttggtt ggtttgttct 840
324 tctatactct atatgctggc tttttggggt ggcaatggcg ccgagttagg actattcaaa 900
326 atgatattaa tgagctcaag aaacaactca aacctgcacc ggtcgcccct gatggtaaag 960
328 cacttgaaac ttcaccgcca tcacctgttg aacttcaaat ccagaaactt actgaggaga 1020
330 ggaaagagct tatcaaagg tcatcacagg ataaacactt taatgctgga tccatacttc 1080
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340 gtatggaaag tgtttgagtt cacaaaatgg ccttgaatgt atgattctca tatgtaagta 1380
342 agttcccagg tattttactt tcaaatcagt atttggcaat atcaataaat gcaaaaattg 1440
344 ctattctgca ttttcaaaaa aaaaaaaaaa aaaaaaaaaa aa 1482

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350 <213> ORGANISM: Medicago sativa

352 <220> FEATURE:

353 <223> OTHER INFORMATION: Nucleotide sequence of genomic H12

355 <400> SEQUENCE: 9

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360 aatttattat taaaattcac acttagatgg cctaaaaatt aacacttatt ttaacaatt      180
362 caaataaaaat atacgacgaa atgagtgtaa tttagttggt taagcatcgt caaagcttgg      240
364 agagaaaagat catagtttga tctttgaaaa ctatactatt gaaaagggtg aagatatcta      300
366 acctccaaca aaatttattt gatagtcgat tcaaattatc aaaatttggg aaatattttg      360
368 taaattgtta agttgggaaa aatatgttaa ttttcaaatt accatttgca catttttcta      420
370 atctcaaadc acatttaagg gatgttgact actttcggtt tgtacaaatc tttacaattt      480
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384 attttcctaa accataccgg tttttccctc ctttatataa ccatcctctc cctcttctc      900
386 taaccaagct tcattcaact cttcaacaca tatcagaaac agaaaaaga agcaaaacat      960
388 tccaagaatt taacaatggc aaccaacgaa gatcaaaagc aaactgaatc tggaagacat     1020
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414 tcatgtaatg cactgctcca cattattgat cattattgtg tggaagctac agagcattta     1800
416 aaagtcttca agccttcttg tcttttgtaa ttttcttca acatatttgt ggttgtaatt     1860
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423 <212> TYPE: PRT

424 <213> ORGANISM: Medicago sativa

426 <220> FEATURE:

427 <223> OTHER INFORMATION: Amino acid sequence encoded by H7 coding region

429 <400> SEQUENCE: 10

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432 1          5          10          15
435 Ala Lys Leu Tyr Lys Ala Leu Ala Lys Asp Ala Asp Glu Ile Val Pro

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 12
Seq#:8; N Pos. 12
Seq#:12; N Pos. 43,44

VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Replaced Current Application Number

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date —

L:164 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0

L:296 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0

L:560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0